SEQUENCE LISTING

(1) GENERAL INFORMATION:

(i) APPLICANT: Warmke, Jeffrey W.
Van Der Ploeg, Leonardus

(ii) TITLE OF INVENTION: PROCESS FOR FUNCTIONAL EXPRESSION OF THE PARA SODIUM CHANNEL

(iii) NUMBER OF SEQUENCES: 7

(iv) CORRESPONDENCE ADDRESS:
(A) ADDRESSEE: Jack L. Tribble
(B) STREET: P.O. Box 2000, 126 E. Lincoln Avenue
(C) CITY: Rahway
(D) STATE: New Jersey
(E) COUNTRY: USA
(F) ZIP: 07065-0907

(v) COMPUTER READABLE FORM:
(A) MEDIUM TYPE: Floppy disk
(B) COMPUTER: IBM PC compatible
(C) OPERATING SYSTEM: PC-DOS/MS-DOS
(D) SOFTWARE: PatentIn Release #1.0, Version #1.25

(vi) CURRENT APPLICATION DATA:
(A) APPLICATION NUMBER:
(B) FILING DATE:
(C) CLASSIFICATION:

(viii) ATTORNEY/AGENT INFORMATION:
(A) NAME: Tribble, Jack L.
(B) REGISTRATION NUMBER: 32,633
(C) REFERENCE/DOCKET NUMBER: 19338DA

(ix) TELECOMMUNICATION INFORMATION:
(A) TELEPHONE: (908) 594-5321
(B) TELEFAX: (908) 594-4720

(2) INFORMATION FOR SEQ ID NO:1:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 33 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

GACTCTAGAC GTTGGCCGCA TAGACAATGA CAG 33

(2) INFORMATION FOR SEQ ID NO:2:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 21 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

AAGAGCTCGA CGAAGGGATC G 21

(2) INFORMATION FOR SEQ ID NO:3:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 24 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:

TCTTCGATCC CTTCGTCGAG CTCT 24

(2) INFORMATION FOR SEQ ID NO:4:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 21 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:

AAAGGATCCA AATATGATGA A 21

(2) INFORMATION FOR SEQ ID NO:5:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 25 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:

TTTGGATCCT TTTTCACACT CAATC 25

(2) INFORMATION FOR SEQ ID NO:6:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 32 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:

GACTCTAGAG CTAATACTCG CGTGCATCTT GG 32

(2) INFORMATION FOR SEQ ID NO:7:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 6513 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:

TCTAGACGTT GGCCGCATAG ACAATGACAG AAGATTCCGA CTCGATATCT GAGGAAGAAC 60

GCAGTTTGTT CCGTCCCTTT ACCCGCGAAT CATTGGTGCA AATCGAACAA CGCATTGCCG 120

CTGAACATGA AAAGCAGAAG GAGCTGGAAA GAAAGAGAGC CGAGGGAGAG GTGCCGCGAT 180

ATGGTCGCAA GAAAAAACAA AAAGAAATCC GATATGATGA CGAGGACGAG GATGAAGGTC 240

CACAACCGGA TCCTACACTT GAACAGGGTG TGCCAATACC TGTTCGATTG CAGGGCAGCT 300

TCCCGCCGGA ATTGGCCTCC ACTCCTCTCG AGGATATCGA TCCCTACTAC AGCAATGTAC 360

TGACATTCGT AGTTGTAAGC AAAGGAAAAG ATATTTTTCG CTTTTCTGCA TCAAAAGCAA 420

TGTGGATGCT CGATCCATTC AATCCGATAC GTCGTGTGGC CATTTACATT CTAGTGCATC 480

CATTATTTTC CCTATTCATC ATCACCACAA TTCTCGTCAA CTGCATCCTG ATGATAATGC 540

CGACAACGCC CACGGTTGAG TCCACTGAGG TGATATTCAC CGGAATCTAC ACATTTGAAT 600

CAGCTGTTAA AGTGATGGCA CGAGGTTTCA TTTTATGCCC GTTTACGTAT CTTAGAGATG 660

CATGGAATTG GCTGGACTTC GTAGTAATAG CTTTAGCTTA TGTGACCATG GGTATAGATT 720

TAGGTAATCT AGCAGCCCTG CGAACGTTTA GGGTGCTGCG AGCGCTTAAA ACCGTAGCCA 780

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ATATGGGCGT GCTCACCGAG AAGTGCATCA AGAAGTTCCC GCTGGACGGT TCCTGGGGCA 960

ATCTGACCGA CGAGAACTGG GACTATCACA ATCGCAATAG CTCCAATTGG TATTCCGAGG 1020

ACGAGGGCAT CTCATTTCCG TTATGCGGCA ATATATCCGG TGCGGGGCAA TGCGACGACG 1080

ATTACGTGTG CCTGCAGGGG TTTGGTCCGA ATCCGAATTA TGGCTACACC AGCTTCGATT 1140

CGTTCGGATG GGCTTTCCTG TCCGCCTTCC GGCTGATGAC ACAGGACTTC TGGGAGGATC 1200

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ATGTGTTTTG TGTGTGGGAC TGTTGCTGGG TTTGGTTGAA ATTTCAGGAG TGGGTATCGC 2460

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CGCTATTGGA ACTGGGACTC GAGGGTGTCC AGGGTCTGTC CGTATTGCGT TCCTTTCGAT 2760

TGCTGCGTGT ATTCAAACTG GCCAAGTCTT GGCCCACACT TAATTTACTC ATTTCGATTA 2820

TGGGACGCAC CATGGGCGCT TTGGGTAATC TGACATTTGT ACTTTGCATT ATCATCTTCA 2880

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TAAGTGATCA ACCATCAGGT GAGAGGACCA ACCAGATCAG TTGGATTTGG AGCGAAGAGC 3360

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CCGCCGTTCT CGTGGAGAGC GACGGGTTCG TGACGAAGAA CGGCCACAAG GTGGTCATCC 6420

ACTCGCGATC GCCGAGCATC ACGTCGCGCA CGGCGGATGT CTGAGCCAGG CCTCGCCCCC 6480

CCCTCCAAGA TGCACGCGAG TATTAGCTCT AGA 6513